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DESIGNATED/ELECTED OFFICE (DO/I				U.S. APPLICATION NO. (If known, see 37 CFR 1.5
	CONCERNING A FILIN			10/009709
INTERN	ATIONAL APPLICATION NO.	INTERNATIONAL FILI	NG DATE	PRIORITY DATE CLAIMED
	SE00/00930	11 May 2000		17 May 1999
TITLEO	FINVENTION	PREVENTION OF DI	ENTAL CARI	ES
	ANT(S) FOR DO/EO/US	Nicklas STROMBE		
Applican	t herewith submits to the United Sta	ates Designated/Elected Off	ice (DO/EO/US)	the following items and other information:
1. 🗶 T	his is a FIRST submission of items	concerning a filing under 3	5 U.S.C. 371.	
2. T	his is a SECOND or SUBSEQUE	NT submission of items con	cerning a filing u	
3. 🔲 T	This is an express request to begin natems (5), (6), (9) and (21) indicated	ational examination procedu below.	ıres (35 U.S.C. 3	71(f)). The submission must include
4. 🔀 T	he US has been elected by the expi	ration of 19 months from th		article 31).
	copy of the International Applicat			nal Rureau)
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b		itted under 35 U.S.C. 154(d)(4).	•
	Amendments to the claims of the Int			(35 U.S.C. 371(c)(3))
a		ed only if not communicate		
Ъ	, <u>—</u>	by the International Bureau.		
С		ever, the time limit for making		ents has NOT expired.
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	An English language translation of t		s under PCT Arti	icle 19 (35 U.S.C. 371 (c)(3)).
9. k A	An oath or declaration of the invento	or(s) (35 U.S.C. 371(c)(4)).	- unsign	ned
	An English lanugage translation of t Article 36 (35 U.S.C. 371(c)(5)).	he annexes of the Internatio	nal Preliminary E	Examination Report under PCT
Items	s 11 to 20 below concern documen	t(s) or information include	e d:	
11.	An Information Disclosure Statem			
12. 🔲	An assignment document for reco	rding. A separate cover she		with 37 CFR 3.28 and 3.31 is included.
13. 🗷	A FIRST preliminary amendment.		EXPRE	ESS MAIL CERTIFICATE
14. 🔲	A SECOND or SUBSEQUENT pr	reliminary amendment.	I hereb	y certify that this correspondence is being
15.	A substitute specification.		deposited with t	the United States Postal Service as Express e to Addresses (mail label
16.	A change of power of attorney and	d/or address letter.	EL3346171	.04US in an envelope addressed to
17. 🔀	A computer-readable form of the s	sequence listing in accorda:	Asst. Commission Novem	oner for Patents, Washington, D.C. 20231. aber 16, 2001
18. 🔲	A second copy of the published in	ternational application und	Dorot	hy Jenkins
19. 🔲	A second copy of the English lang	guage translation of the inte	Name (of Person Mailing Correspondence
20. 🗶	Other items or information: PEFS Print form.		Xlor	Signature /
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21. X The following fees are submitted	CALCULATIONS PTO USE ONLY					
BASIC NATIONAL FEE (37 CFR 1.492						
Neither international preliminary exami	nation fee (37 CFR 1.482)					
nor international search fee (37 CFR 1.4 and International Search Report not pre	oared by the EPO or JPO	\$1040.00				
International preliminary examination for USPTO but International Search Report	ee (37 CFR 1.482) not paid to	\$890.00		,		
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but international search fee (37 CFR 1.4	45(a)(2)) paid to USPTO	\$740.00				
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but all claims did not satisfy provisions	of PCT Article 33(1)-(4)	\$/10.00				
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and all claims satisfied provisions of PC ENTER APPROPRIA			\$11040.00			
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Surcharge of \$130.00 for furnishing the commonths from the earliest claimed priority	date (37 CFR 1.492(e)).	20 30	\$ 1040.00			
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MULTIPLE DEPENDENT CLAIM(S) (i	f applicable) L OF ABOVE CALCU	<u> </u>	\$ 1040.00			
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	TOTAL NATIO	ONAL FEE =	\$: 1040.00			
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OSTROLENK, FABER, GERB & 1180 Avenue of the Ame						
New York, NY 10036-8			rd A. Meilma	n		
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19 NOV 2001 P/2432-44

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Patent Application of					
Nicklas Strömberg et al.	e: No	November 16, 2001			
Serial No	up A	Art Unit:			
Filed:	Exa	minei	. :		
For: PREVENTION OF DENTAL CARIES					
Asst. Commissioner for Patents					
Washington, D.C. 20231					
AMENDMENT/SUBM	ISSI	ON			
Prior to examination, please amend the application a	as foll	lows.			
FEE CALCULATION					
Any additional fee required has been calculated as for	ollow	s:			
If checked, "Small Entity" status is c	laime	d.			
NO. CLAIMS HIGHEST NO.					
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INDEP. 1 MINUS 3 **=	0	X	(\$42 SE or \$84)	\$	
FIRST PRESENTATION OF MULTIPLE DEPENDENT CLAIM		X	(\$140 SE or \$280)	\$	
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enclosed or if any additional fee during the prosecution of t	his ap	plica	tion is not paid, th	e Pat	tent
Office is authorized to charge the underpayment to Deposit	Acc	ount]	No. 15-0700.		

CONTINGENT EXTENSION REQUEST

If this communication is filed after the shortened statutory time period had elapsed and no separate Petition is enclosed, the Commissioner of Patents and Trademarks is petitioned, under 37 C.F.R. §1.136(a), to extend the time for filing a response to the outstanding Office Action by the number of months which will avoid abandonment under 37 C.F.R. §1 135. The fee under 37 C.F.R. § 1.17 should be charged to our Deposit Account No. 15-0700.

AMENDMENTS

X If checked, amendment(s) to the specification and/or claims are submitted herewith.

Claims:

Please amend claims 5, 6 and 8 and add new claims 9-17 pursuant to 37 C.F.R. § 1.121(c)(i) as set forth in the "clean" version attached hereto as Appendix A. Entry is respectfully requested. A version with markings to show the changes made pursuant to 37 C.F.R. § 1.121(c)(ii) is attached hereto as Appendix B.

REMARKS/ARGUMENT

This Preliminary Amendment is being submitted to change the multiple dependent claims to single dependent claims in order to eliminate the improper multiple dependent claims and to reduce the government filing fee.

EXPRESS MAIL CERTIFICATE

I hereby certify that this correspondence is being deposited with the United States Postal Service as Express Mail to Addressee (mail label # EL534617104US) in an envelope addressed to: Asst. Commissioner for Patents, Washington, D.C. 20231, on November 16, 2001

Dorothy Jenkins

Name of Person Mailing Correspondence

November 16, 2001

Date of Signature

Respectfully submitted,

Edward A. Meilman

Registration No.:24,735

OSTROLENK, FABER, GERB & SOFFEN, LLP

1180 Avenue of the Americas

New York, New York 10036-8403

Telephone: (212) 382-0700

APPENDIX A

"CLEAN" VERSION OF EACH PARAGRAPH/SECTION/CLAIM 37 C.F.R. § 1.121(b)(ii) AND (c)(i)

CLAIMS (with indication of amended or new):

- 5. (Amended) A method of preventing dental caries comprising the oral administration of a prevention-effective amount of the peptide of claim 1.
- 6. (Amended) A composition for preventing dental caries comprising a prevention-effective amount of the oligopeptide of claim 1.
- 8. (Amended) The manufacture of a medicament for prevention of dental caries comprising a prevention-effective amount of the oligopeptide of claim 1.
- 9. (NEW) A method of preventing dental caries comprising the oral administration of a prevention-effective amount of the peptide of claim 2.
- 10 (NEW) A method of preventing dental caries comprising the oral administration of a prevention-effective amount of the peptide of claim 3.
- 11. (NEW) A method of preventing dental caries comprising the oral administration of a prevention-effective amount of the peptide of claim 4.
- 12. (NEW) A composition for preventing dental caries comprising a prevention-effective amount of the oligopeptide of claim 2.
- (NEW) A composition for preventing dental caries comprising a prevention-effective amount of the oligopeptide of claim 3.
- 14. (NEW) A composition for preventing dental caries comprising a prevention-effective amount of the oligopeptide of claim 4.

- 15. (NEW) The manufacture of a medicament for prevention of dental caries comprising a prevention-effective amount of the oligopeptide of claim 2.
- 16. (NEW) The manufacture of a medicament for prevention of dental caries comprising a prevention-effective amount of the oligopeptide of claim 3.
- 17. (NEW) The manufacture of a medicament for prevention of dental caries comprising a prevention-effective amount of the oligopeptide of claim 4.

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APPENDIX B

VERSION WITH MARKINGS TO SHOW CHANGES MADE 37 C.F.R. § 1.121(b)(iii) AND (c)(ii)

CLAIMS:

- 5. (Amended) A method of preventing dental caries comprising the oral administration of a prevention-effective amount of the peptide of [any of claims] <u>claim</u> 1[-4].
- 6. (Amended) A composition for preventing dental caries comprising a prevention-effective amount of the oligopeptide of [any of claims] <u>claim</u> 1[-4].
- 8. (Amended) The manufacture of a medicament for prevention of dental caries comprising a prevention-effective amount of the oligopeptide of [any of claims] claim 1[-4].

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531 Rec'd PCT/TT 19 NOV 2001

Date: November 16, 2001

Group Art Unit: --

Examiner: --

P/2432-44

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Patent Application of

Nicklas Strömberg et al.

Serial No.: Not Yet Known

Filed: Herewith

For: PREVENTION OF DENTAL CARIES

Assistant Commissioner for Patents Box Patent Application

Washington, D.C. 20231

STATEMENT IN ACCORDANCE WITH 37 C.F.R. §1.821(f)

Sir:

Pursuant to 37 C.F.R. §1.821(f), I hereby certify that the contents of the sequence listings enclosed in the above-identified patent application papers and the contents of the computer readable form of the sequence listings are the same.

I hereby declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any parent issued thereon.

Respectfully submitted,

Charles C. Achkar

Registration No.: 43,311

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New York, New York 10036-8403

Telephone: (212) 382-0700

PREVENTION OF DENTAL CARIES

FIELD OF THE INVENTION

5 The present invention relates to a means for preventing dental caries, a corresponding method, and compositions incorporating the means.

BACKGROUND OF THE INVENTION

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Proline-rich proteins (PRPs) comprise about 70% of the proteins in saliva. They can be divided into acidic, basic, and glycosylated PRPs encoded by six clustered genes on chromosome 12p13.2. PRPs are potential determinants of host susceptibility to dental caries. Large multifunctional 150-15 residue PRPs (Lamkin M S and Oppenheim F G, Crit Rev Oral Biol Rev 4 (1993) 251-9) are known to adsorb to hydroxyapatite surfaces and inhibit calcium phosphate precipitation, modulate hydroxyapatite crystal formation as well as mediate adhesion of commensal Actinomyces and 20 Streptococcus species to tooth surfaces. In addition, they bind to and inactivate ingested plant polyphenols (tannins). The proline-poor N-terminal 30 residue domain confers hydroxyapatite- and calcium-binding, while the proline-rich middle/C-terminal domain binds bacteria via the ProGln-25 terminus and tannins via proline-rich repeats.

The polymorphism of acidic PRPs involves allelic and post-translational variants; several common allelic (large PRP-1, PRP-2, Db-s, PIF-s and Pa) and post-translational (small PRP-3, PRP-4, Db-f and PIF-f) variants have been described. In regard of biological properties the acidic PRP variants differ somewhat, and large and small acidic PRPs largely,

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between each other. The small 106 residue acidic PRPs resulting from proteolytic cleavage at Arg106-Gly107 display poor bacterial adhesion activity but high affinity for hydroxyapatite surfaces. Both large and small acidic PRPs are secreted from the acinus cells. After secretion, the acidic PRPs are rapidly enriched on tooth surfaces and degraded as a consequence of bacterial proteolysis. Also, multiple phosphorylated peptides reminiscent of PRPs have been identified in saliva and found to possess increased affinity for hydroxyapatite surfaces. Some studies (Kousvelari E E et al., J Dent Res 59 (1980) 1430-8), though not others (Mandel I D & Bennick A, J Dent Res 62 (1980) 943-5), have associated this proteolytic activity to gingivitis and rate of plaque formation.

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Arg catabolism to ammonia has been suggested to be a characteristic of streptococcal biotypes with tooth protective properties (Andersson C et al. Infect Immun 43 (1984) 555-60; Rogers AH, Aust Dent J 35 (1990) 468-71. Wijeyeweera R L& Kleinberg, Arch Oral Biol 34 (1989) 55-64; 20 ibid. 43-53). Actually, Arg can control the ecological relationship between S. sanguis and S. mutans (van der Hoeven J S et al., J Dent Res 63 (1984) 389-92. Rogers A H et al., Oral Microbiol Immunol 2 (1987) 172-82), and cariessusceptible and caries-resistant subjects differ in ability 25 to raise pH after an acidification (Abelson D C & Mandel I D, J Dent Res 60 (1981) 1634-8; Kleinberg I et al., Proc. "Microbial aspects of dental caries". Sp. Supp. Microbiol. Abstracts. IRL, Washington, D.C., 1976, 433-464 pp).

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Arg catabolism is present in *S. sanguis*, *S. gordonii*, *S. anginosus* and *S. mitis*, while it is absent in *S. oralis* and cariogenic *S. mutans* and *S. sobrinus* (Kilian M et al., Int J

Syst Bacteriol 39 (1989) 471-484). Arg-containing peptides, notably sialin (Coulter W A et al., Biochem Soc Trans 18 (1990) 337-8) may act as a local pH-buffering agents by bacterial catabolism of Arg to ammonia via the arginine deaminase pathway (Floderus E et al., APMIS 98 (1990) 1045-52).

Degradation of acidic PRPs releases oligopeptides which are transported intra-cellularly and metabolized for nutritional reasons (Kunji E R et al., Antonie Van Leeuwenhoek 70 (1996) 10 187-221). Gram-negative bacteria colonizing the gingival pocket, such as Porphyromonas gingivalis, express a broad range of proteases in utilizing proteins as a major source of energy. In contrast, commensal streptococci and actinomycetes, which dominate on oral mucosal and tooth 15 surfaces, express multiple glycosidases in utilizing carbohydrates as a major energy source. Nevertheless, protease and peptidase activities, such as a trypsin-like serine endoprotease in S. oralis, S-IgA1 proteases in S. 20 sanguis and S. oralis and yet uncharacterized proteolytic activity, are present in Streptococcus and Actinomyces species. However, little is known about the degradation of acidic PRPs by these commensal and early colonizing bacteria.

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While a vast amount of knowledge about saliva constitutents and their possible role in processes leading to dental caries has been gathered, that knowledge up to know resulted in only few practically applicable propositions of how to efficiently prevent dental caries.

OBJECTS OF THE INVENTION

It is an object of the invention to provide a means for protecting dental surfaces against caries.

It is another object of the invention to provide a corresponding method.

It is a further object of the invention to provide a composition for protecting dental surfaces against caries.

10 Still other objects of the invention will become apparent from the following description of the invention and preferred embodiments thereof, as well as from the appended claims.

15 DESCRIPTION OF THE INVENTION

The present invention is based on the insight that commensal Actinomyces and Streptococcus species transform acidic PRPs to small-size peptides, such as pentapeptides. These small-size peptides are transformed into ammonia by the action of certain oral bacteria. The ammonia thus formed protects raises the pH at the dental surface and therby protects the surface against caries.

In this transformation Arg-containing oligopeptides are released and catabolized to ammonia at or near the dental surface. Ammonia increases the pH at the dental surface and thereby counteracts caries which is promoted by a low pH in the oral cavity.

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The present invention is also based on the insight that the aforementioned transformation gives rise to peptides with terminal proline or proline-glutamine entities having

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affinity to dental surfaces and thus competitively protecting them.

Degradation of acidic PRPs by Streptococcus and Actinomyces species comprises the following features:

- (i) step-by-step the large 150 residue PRP-1 protein is degraded into intermediate size peptides and finally into smaller oligopeptides and amino acids. Cell-bound endoproteases with broad substrate specificities are
- involved in the initial stages of this degradation;

 (ii) cleavage of acidic PRPs occurs preferentially in the middle/C-terminal proline-rich 96-140 region; the 106 residue PRP-3 is relatively resistant to cleavage, PRP-3-like peptides being generated by all strains and a series of fragments in region 96-140. Both endoprotease specificity per se and intrinsic properties of acidic PRPs, such as the extended structure of proline-rich stretches, explains this regional preference of cleavage;
- (iii) degradation of PRP-1 releases oligopeptides both
 immediately and after prolonged degradation. The immediate release of an Arg106Gly107Arg108Pro109Gln110 pentapeptide is indicated by the N-terminal 105 residue Pyr1-Pro104Pro105 and the C-terminal 40 residue Gly111-Pro149Gln150 peptides initially generated by S. gordonii strain SK12.

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Mass spectroscopy identified a N-terminal pyroglutamatic acid (Pyr) post-translational modification of acidic PRPs that blocks sequencing by Edman degradation. The peptide structures derived from degradation of the 150 residue PRP-1 by *S. gordonii* strain SK12 indicate:

(i) the instant release of an Arg106Gly107Arg108Pro109Gln110 pentapeptide together with a N-terminal 105 residue Pyr1-

Pro104Pro105 and a C-terminal 40 residue Gly111-Pro149Gln150
peptide;

- (ii) the generation of multiple middle/C-terminal peptides with ProGln-termini;
- 5 (iii) the cleavage at peptide bonds formed at Pro or Gln residues: Pro95-Pro96, Pro104-Pro105, Pro105-Arg106, Gln110-Gly111, Gln131-Gly132, Pro135-Gln136, Gln136-Gly137, and Gln141-Gly142.
- Degradation of acidic PRPs by *S. gordonii* SK12 initially generates a pentapeptide, Arg106Gly107Arg108Pro109Gln110, containing two of the five Arg residues in PRP-1.
- The N-terminal Glu1-Pro104Pro105 peptide derived from PRP-1
 by S. gordonii SK12 lacks the bacterial adhesion-promoting
 ProGln-terminus, but contains the hydroxyapatite binding
 domain. Since small phosphorylated proline-rich peptides and
 PRP-3 display increased affinity for hydroxyapatite surfaces
 (Moreno E C et al., J Biol Chem 257 (1982) 2981-9;
- Madapallimattam G & Bennick A, Biochem J 270 (1990) 297-304;
 Lamkin M S et al., J Dent Res 75 (1996) 803-8), such Nterminal peptides can affect bacterial adhesion by occupying
 hydroxyapatite sites or altering the pellicle mosaic of
 binding structures. Proteolytic cleavage is known to unmask
- hidden binding structures (Gibbons R J et al., Arch Oral Biol 35 Suppl (1990) 107s-114s), and strains of Actinomyces species recognize diverse sites in acidic PRPs and statherin (Hallberg K et al., Oral Microbiol Immunol 13 (1998) 327-336; Li T et al., Infect Immun 67 (1999) 2053-9).
- 30 Furthermore, acidic PRP-degradation generated multiple middle/C-terminal peptides with adhesion-active ProGln-termini, potentially inhibiting the adhesion of oral pathogens binding to acidic PRPs and statherin, i. e. S.

mutans (Gibbons R J & Hay D I, J Dent Res 68 (1989) 1303-7), P. gingivalis (Amano A et al., Infect Immun 62 (1994) 3372-80) and Fusobacterium nucleatum (Xie H et al., Oral Microbiol Immunol 6 (1991) 257-63). Commensal and pathogenic species display high and low avidity binding to acidic PRPs, respectively, and may respond differently to ProGlnterminating peptides. Alternatively, both the N-terminal and middle/C-terminal peptides could act as inhibitors or retarders of plaque formation in a broader sense.

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While the inventors believe these explanations to be true they are of a hypothetical nature and should not be understood so as to limit the scope of the invention which is only determined by the appended claims.

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According to the invention is disclosed an oligopeptide protecting against dental caries comprising two arginine residues selected from the group consisting of pentapeptide, hexapeptide, heptapeptide, octapeptide, nonapeptide and decapeptide. Particularly preferred is the pentapeptide ArgGlyArgProGln.

Also preferred is a penta- to decapeptide comprised by the sequence of amino acid 99 to amino acid 115 of the 150 residue PRP-1 protein:

GlyGlyHisProArgProProArgGlyArgProGlnGlyProProGlnGln, with the provisio that it contains two or more Arg.

Also preferred are the following peptides:

ArgGlyArgProGln (residues 106-110);

ArgGlyArgProGlnGly (residues 106-111);

ArgGlyArgProGlnGlyPro (residues 106-112);

ArgGlyArgProGlnGlyProPro (residues 106-113);

ArgGlyArgProGlnGlyProProGln (residues 106-114);
ArgGlyArgProGlnGlyProProGlnGln (residues 106-115);
GlyGlyHisProArgProProArgGlyArg (residues 99-108);
GlyHisProArgProProArgGlyArg (residues 100-108);
HisProArgProProArgGlyArg (residues 101-108);
ProArgProProArgGlyArg (residues 102-108);
ArgProProArgGlyArg (residues 103-108);
ProProArgGlyArg (residues 104-108).

10 The peptides of the invention can be easily synthesized by solid phase or liquid phase methods known in the art.

According to the invention is disclosed a method of preventing dental caries comprising the oral administration of a prevention-effective amount of an oligopeptide comprising two arginine residues selected from the group consisting of pentapeptide, hexapeptide, heptapeptide, octapeptide, nonapeptide and decapeptide. Particularly preferred is the pentapeptide ArgGlyArgProGln for use in the method of preventing dental caries.

Also preferred for use in the method of preventing dental caries is a penta- to decapeptide comprised by the sequence of amino acid 99 to amino acid 115 of the 150 residue PRP-1 protein:GlyGlyHisProArgProProArgGlyArgProGlnGlyProProGlnGln, with the provisio that it contains two or more Arg.

Also preferred for use of preventing dental caries are the following peptides:

ArgGlyArgProGln (residues 106-110);
ArgGlyArgProGlnGly (residues 106-111);
ArgGlyArgProGlnGlyPro (residues 106-112);
ArgGlyArgProGlnGlyProPro (residues 106-113);

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ArgGlyArgProGlnGlyProProGln (residues 106-114);
ArgGlyArgProGlnGlyProProGlnGln (residues 106-115);
GlyGlyHisProArgProProArgGlyArg (residues 99-108);
GlyHisProArgProProArgGlyArg (residues 100-108);
HisProArgProProArgGlyArg (residues 101-108);
ProArgProProArgGlyArg (residues 102-108);
ArgProProArgGlyArg (residues 103-108);
ProProArgGlyArg (residues 104-108).
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- 10 According to the invention is disclosed a composition for preventing dental caries comprising a prevention-effective amount of an oligopeptide comprising two arginine residues selected from the group consisting of pentapeptide, hexapeptide, heptapeptide, octapeptide, nonapeptide and decapeptide, and a suitable carrier. Particularly preferred is the pentapeptide ArgGlyArgProGln. Suitable carriers include state-of-the-art toothpaste and mouthwash compositions but also chewing-gums, lozenges, and the like.
- Also preferred for use in the composition for preventing dental caries is a penta- to decapeptide comprised by the sequence of amino acid 99 to amino acid 115 of the 150 residue PRP-1 protein:

 GlyGlyHisProArgProProArgGlyArgProGlnGlyProProGlnGln,
 with the provisio that it contains two or more Arg.

Also preferred for use in the composition for preventing dental caries are the following peptides:

ArgGlyArgProGln (residues 106-110);

ArgGlyArgProGlnGly (residues 106-111);
ArgGlyArgProGlnGlyPro (residues 106-112);
ArgGlyArgProGlnGlyProPro (residues 106-113);
ArgGlyArgProGlnGlyProProGln (residues 106-114);
ArgGlyArgProGlnGlyProProGlnGln (residues 106-115);
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GlyGlyHisProArgProProArgGlyArg (residues 99-108);
GlyHisProArgProProArgGlyArg (residues 100-108);
HisProArgProProArgGlyArg (residues 101-108);
ProArgProProArgGlyArg (residues 102-108);
ArgProProArgGlyArg (residues 103-108);
ProProArgGlyArg (residues 104-108).

Since the caries-protecting peptides according to the invention are peptides produced by the human body or correspond to sequences thereof they are well tolerated. Optimal caries-protecting amounts of these peptides can be easily determined by experiment because the absence of important toxicity restraints. Single doses from 0.1 mg to 1 g will be appropriate for most applications.

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The invention will be better understood by reference to a drawing and the following description of preferred embodiments thereof.

20 DESCRIPTION OF THE DRAWING

- Fig. 1 illustrates the native alkaline PAGE patterns of N-terminal acidic peptides from degradation of acidic PRPs (PRP-1 and PRP-3) by strains of commensal Streptococcus and Actinomyces species;
- Fig. 2 illustrates the gel filtration patterns of fragments formed by degradation of PRP-1 by strains representative for each of the four degradation patterns identified among streptococci and actinomycetes;
- Fig. 3a illustrates the gel filtration patterns of peptide fragments from the degradation of PRP-1 by S.

 gordonii strain SK12 for different time periods;

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Fig. 3b illustrates the identification by mass spectrometry of fragments separated using either gel filtration (peaks 1-4, cf. Fig. 3a) or peptide gel filtration (peak 5).

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DESCRIPTION OF PREFERRED EMBODIMENTS

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EXAMPLE 1. Isolation of acidic PRPs. Parotid saliva was collected from three subjects homozygous for PRP-1 and PIF-s using Lashley cups and mild acidic lozenge stimulation. After pooling of the salivas and dilution (1:1 by volume) with 50 mmol/L Tris-HCl, 25 mmol/L NaCl, pH 8.0 (Tris-HCl 15 buffer), the saliva sample was subjected to DEAE-Sephacel column chromatography (15 x 1.6 cm, Pharmacia, Uppsala, Sweden) using a linear gradient of 25 to 1000 mmol/L NaCl in Tris-HCl buffer. The protein fraction containing the acidic PRPs (PRP-1/PIF-s and PRP-3/PIF-f) was concentrated via 20 ultrafiltration using a Centriprep 10 concentrator (Amicon Inc., Beverly, MA), and subjected to gel filtration (HiLoad™ 26/60 Superdex S-200 Prep grade column, Pharmacia) in 20 mmol/L Tris-HCl, 500 mmol/L NaCl, pH 8.0. The resolved protein fractions (PRP-1/PIF-s and PRP-3/PIF-f) were 25 dialyzed against Tris-HCl buffer (Spectra/Pore membrane No. 4, Spectrum Medical Industries, Inc., Houston, TX). Each protein fraction (PRP-1/PIF-s and PRP-3/PIF-f) was finally purified on a Macroprep high Q column (15 x 1.6 cm, Bio-Rad, Hercules, CA) using a linear gradient of 25 to 1000 mmol/L 30 NaCl in 50 mmol/L Tris-HCl, pH 8.0. The purified proteins were extensively dialyzed against water, lyophilized and stored at -20°C.

EXAMPLE 2. Bacterial strains and culturing. The origins and species designations of strains of Actinomyces naeslundii, Actinomyces odontolyticus, A. naeslundii genospecies 1 strain ATCC 12104, genospecies 2 strain T14V, A. 5 odontolyticus strain PK984 and A. viscosus strain 19246 are as previously described (Hallberg K et al, see above; Hallberg K et al., Oral Microbiol Immunol 13 (1989) 188-192; Li et al., 1999, see above). The strains of S. anginosus, S. gordonii, S. mitis, S. oralis and S. sanguis (provided by 10 Dr. M. Kilian, Department of Oral Biology, Royal Dental College, Aarhus, Denmark) have been described (Kilian et al., 1989; Hsu et al., 1994). S. mutans strains Ingbritt, JBP and NCTC 10449, S. sobrinus strain SL-1 and Porphyromonas gingivalis strains W83 and ATCC 33277 were 15 provided by Dr. J. Carlsson, Department of Oral Biology, Umeå University, Sweden. All strains were grown overnight on Columbia-II-agar base plates (Becton Dickinson and Company, Cockeysville, MD), supplemented with 30 ml of a human erythrocyte suspension per liter, at 37°C in an atmosphere 20 with 5 % CO2. Growth of S. gordonii strain SK12 identified two morphological types, one forming flat (strain SK12-I) and the other convex (strain SK12-II) colonies. Strain SK12-I (the flat colony-forming type) was selected for the degradation experiments. Neither whole cell soluble protein 25 patterns, as analyzed by SDS-PAGE, nor acidic PRP cleavage patterns distinguished between the two types. The strains intended for degradation experiments were then grown at 37°C for 18 h in 5 mL trypticase soy bean glucose limiting broth (1.7% peptone, 0.3% soy peptone, 0.15% yeast extract, 12.5 30 mmo1/L glucose, 12.5 mmo1/L NH_4HCO_3 in 1 mo1/L NaH_2PO_4/K_2HPO_4 buffer, pH 7.3) in an atmosphere with 5% CO_2 . The cells were pelleted by centrifugation at 17,000 x g for 5 min, washed

twice in 0.43% NaCl, 0.042% KCl, 0.1% Na₂HPO₄, 0.1% KH₂PO₄,
 1% glycerophosphate disodium salt, 0.024% CaCl₂, and 0.01%
 MgCl₂.H₂O (M-DIL buffer), re-suspended in M-DIL buffer at a
 concentration of 2 x 109 cells/mL. The bacterial cells (and
 cell-free supernatant after pelleting of bacteria by
 centrifugation) were kept on ice prior to degradation
 experiments. The strains intended for adhesion tests were
 metabolically labeled by adding 35S-methionine (200 *Ci,
 Tran 35S-Label, ICN Pharmaceuticals Inc., Irvine, CA) to
 bacteria suspended in 100 μL 10 mmol/L Na₂HPO₄, NaH₂PO₄,
 0.137 M NaCl, pH 7.2, (PBS) prior to growth on Columbia-II agar plates.

EXAMPLE 3. Degradation of acidic PRPs. Equal volumes (200 μL) of purified protein (0.6 mg/mL) and bacteria (2 x 109 cells/mL), both dissolved in M-DIL buffer, were mixed and incubated at 37°C for 15 min, 4 h, 20 h and 1 week. After pelleting of the bacteria by centrifugation at 17,000 x g for 10 min, the supernatants were aliquoted, lyophilized and stored at -80°C prior to analysis. In some experiments, cell free enzyme supernatants and parotid saliva from defined acidic PRP phenotypes were used following the same protocol. The saliva was collected as described above and sterilized by filtering (0.20 μm Minisart filter, Sartorius, Göttingen, Germany).

EXAMPLE 4. Native alkaline PAGE. Native alkaline polyacrylamide gel electrophoresis was performed essentially as described (Azen and Yu, 1984). The lyophilized supernatants were dissolved in 50 μ L sample buffer containing 1% glycine, 10% glycerol and 0.025% bromophenol blue and centrifuged at 17,000 x g for 10 min. The electrophoresis was performed using precast Tris-glycine

7.5% resolving gels with 4% stacking gels (Bio-Rad) at 100 V for 20 min followed by 175 V for 1 h 30 min in a buffer containing 0.038 M Tris-glycine, pH 8.4. The gels were stained with 0.1% Coomassie Blue R (Serva Feinbiochemica, Heidelberg, Germany) in 20% trichloroacetic acid (Merck, Darmstadt, Germany) for 16 h, destained in 2% acetic acid for 16 h and finally soaked in water for 2 h.

EXAMPLE 5. Densitometry. The degree of acidic PRP

degradation was quantified by densitometry of native
alkaline PAGE gels using a Model GS-700 Imaging densitometer
and the Molecular Analyst Software (Bio-Rad). The degree of
degradation of PRP-1 and PRP-3 was scored from 0 to 5
according to the following criteria: 0 = 0<10%, 1 = 10<20%,

2 = 20<40%, 3 = 40<60%, 4 = 60<80%, 5 = 80<100% reduction of
PRP-1 or PRP-3 at 4 h of incubation. In the case of certain
strains, score 1 denotes the formation after 20 h of
incubation of peptide fragments similar to those generated
after 4 h by the strains with a high PRP degradation rate

(Table 1).

EXAMPLE 6. **Gel filtration.** The lyophilized supernatants obtained by incubation of bacteria with acidic PRPs were dissolved in 200 μL 20 mM Tris-HCl, pH 8.0, containing 0.5 M NaCl and subjected to gel filtration on Superose 12 HR 10/30 or a Superdex Peptide HR10/30 (Pharmacia) columns, equilibrated in 20 mM Tris-HCl, pH 8.0. The flow rate was 0.2 mL/min and the absorbance was monitored at 214 nm. Fractions of 1 mL were collected.

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EXAMPLE 7. Hydroxyapatite assay. Adherence of 35S-methionine labeled bacteria to acidic PRP-1 was measured by the

hydroxyapatite assay as previously described (Carlén et al., 1998).

EXAMPLE 8. Mass spectrometry. All mass spectrometric data were acquired on a hybrid quadrupole time-of-flight (Q-ToF) 5 mass spectrometer (Micromass, Manchester, UK). A Zconfigured nano spray source was used and samples were introduced via gold-coated spraying needles (Protana, Odense, Denmark). Detection was all times in the positive ion mode. Needles were opened manually under a light-10 microscope resulting in an opening of approximately 5 μm . Before analysis, remaining contaminants and salt were removed by applying the samples to nano-columns; Gel-loader tips (Eppendorf, Hamburg, Germany) packed with POROS R3 reverse phase resin (PerSeptive Biosystems, Framingham, MA, 15 USA) to a bed volume of approximately 200 nL. Normally, 5 μL of sample was applied to the resin followed by washing with approximately 30 μ L 0.1% TFA and elution with 60% CH3CN in 1% acetic acid directly into the nano-spray needle.

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- EXAMPLE 9. N-terminal sequence analysis. Edman degradation was performed directly on proteins blotted (Transblot, Bio-Rad) onto PVDF-membranes using an Applied Biosystems 477A pulsed liquid phase sequencer (Foster City, CA) with an online PTH 120A analyzer. Sequence analysis was performed with cycle programs adapted to the reaction cartridges and chemicals from the manufacturer. Yields were calculated from sequenced standard *-lactoglobulin.
- EXAMPLE 10. Substrate specificity. The substrate specificity of bacterial strains was measured as previously described (Erlanger et al., 1961). Briefly, 25 μ L of bacterial suspension (5 x 109 cells/mL in M-DIL buffer) was diluted

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with 25 μ L of 0.1 mol/L Tris-HCl, pH 7.0, followed by addition of 50 μ L of B-Arg-pNA (Sigma, St. Louis, MO) and Z-Gly-Pro-pNA (Bachem, Bulendorf, Switzerland) (8.0 mmol/L in DMSO). After incubation at 37°C for 16 h, the extent of cleavage was measured by the absorbance at 414 nm.

EXAMPLE 11. Degradation of acidic PRPs by commensal Streptococcus and Actinomyces species. Strains of Streptococcus and Actinomyces species were tested for degradation of acidic PRPs (PRP-1 and PRP-3) by co-10 incubation of bacteria and protein followed by native alkaline PAGE (Table 1, Fig. 1). Degradation of PRP-1 occurred with S. oralis (1 of 4 strains), S. mitis (1 of 4 strains), S. gordonii (3 of 4 strains), S. sanguis (4 of 4 strains), S. anginosus (3 of 4 strains) and A. odontolyticus 15 (4 of 7 strains), but not with S. mutans (3 strains), S. sobrinus (1 strain), A. naeslundii genospecies 1 (6 strains) and genospecies 2 (7 strains) or A. viscosus (1 strain). S. gordonii and A. odontolyticus displayed high PRP-1 cleavage rates (scores 4 and 5), while other species had moderate 20 (scores 2 and 3) to low (score 1) cleavage rates. While most species showed high PRP-1 relative to PRP-3 cleavage rates, A. odontolyticus displayed equal PRP-1 and PRP-3 cleavage rates. Degradation of PRP-1 also occurred in cell-free supernatants (Table 1), and S. gordonii strain SK12 (with a 25 high PRP-1 cleavage rate) degraded PRP-1, PRP-2, Db-s, PIF-s and Pa in parotid saliva from defined donors (data not shown).

30 Both streptococci adhering to and those not adhering to PRP1 degraded acidic PRPs (Table 1). While A. naeslundii
genospecies 1 and 2 bound to but did not degrade acidic
PRPs, the opposite was true for A. odontolyticus. In

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addition, both S-IgAl protease positive and negative strains degraded acidic PRPs (Table 1). Thus, no obvious relationship was found between PRP degradation and adhesion to acidic PRPs or S-IgAl protease activity.

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EXAMPLE 12. Time-dependency of acidic PRP degradation. The native alkaline PAGE patterns from degradation of acidic PRPs for different times indicated a time-dependency of degradation for all strains (Fig. 1). The time-dependency of degradation of PRP-1 by S. gordonii strain SK12 was further analyzed by gel filtration of fragments formed after different times of incubation (Fig. 3a). Degradation of PRP-1 (peak 1) for 15 min generated a peptide eluting similar to PRP-3 (peak 2) and later-eluting, smaller peptides (peak 3). Prolonged degradation (20 h) completely converted PRP-1 into the peptide eluting similar to PRP-3 (peak 2) and even later-eluting peptides (peak 4). Even further degradation (1 week) completely transformed all peptides (except for the one eluting similar to PRP-3) into oligopeptides and amino acids (peak 5), as identified using peptide gel filtration (data not shown).

derived peptides. The peptides generated from PRP-1 by S.

gordonii strain SK12 were identified by mass spectrometry of the peptide peaks obtained by gel filtration (Figs 2 and 3).

An N-terminal 105 residue peptide Pyr1-Pro104Pro105 (peak 2) and a C-terminal 40 residue peptide Gly111-Pro149Gln150 (peak 3) were identified after initial (15 min) cleavage of PRP-1 (Fig. 3b). A series of 15-47 residue peptides (peak 4; Pro96-Pro109Gln110, Gly111-Pro130Gln131, Gly111-Pro134Pro135, Gly111-Pro135Gln136, Gly111-Pro140Gln141, Gly111-Pro149Gln150 and Pro104-Pro149Gln150) were identified

after prolonged degradation (20 h). In addition, a pyroglutamatic acid, Pyr, was found at the N-terminal residues of PRP-1 and PRP-3 (Fig. 3b), explaining our difficulties in sequencing of N-terminal fragments by Edman degradation.

EXAMPLE 14. Patterns of acidic PRP degradation. The peptide profiles displayed by native alkaline PAGE distinguished four major degradation patterns, I-IV (Table 1, Fig. 1). While all four patterns contained N-terminal peptides 10 migrating close to PRP-3, the peptide banding patterns were unique to a particular strain or species (Fig. 1, Table 1). Both species specific (A. odontolyticus and S. gordonii) and multiple patterns within a species (S. anginosus and S.sanguis), as well as a sharing of patterns between species, 15 were observed. In addition, gel filtration of the degradation products obtained by a strain representative of each pattern verified somewhat deviating profiles, containing large (70-75 min retention) and intermediate (80-95 min retention) peptides (Fig 1b). To investigate the 20 substrate specificities associated with acidic PRP degradation, strains representative of each degradation pattern were tested for truncation of Pro- and Argcontaining chromogenic substrates. The strains did not cleave Z-Gly-Pro-pNA, which is a substrate for prolyl 25 endoproteases (Blumberg et al., 1980), or B-Arg-pNA, while P. gingivalis strain ATCC 33277 cleaved both substrates.

EXAMPLE 15. Lozenge. A solution of ArgGlyArgProGln

'acetate' was prepared by dissolving ArgGlyArgProGln in
water and adding acetic acid to pH 6.5. The aqueous solution
was freeze-dried and the powder thereby obtained mixed with

150 g of polyyethylene glycol 8000, 150 g of

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microcrystalline cellulose, 600 g of mannitol, 10 g of stearic acid are milled to pass a 40 mesh sieve. The mixture is fed to a tablet press to produce 1 g tablets.

5 EXAMPLE 16. Chewable tablet. 900 g mannitol and 5 g sodium saccharin are screened through a 40-mesh screen and blended thoroughly with 40 g ArgGlyArgProGln acetate prepared as described above. A binder solution of 20 g of acacia and 50 g of gelatin in 500 ml water was prepared separately. The 10 powder was wet granulated using 200 ml of binder solution for 1000 powder. After drying overnight at 75°C the granules were screened through a 12 mesh screen, mixed with 1 g of peppermint oil adsorbed on 3 g of colloidal silica and 25 g magnesium stearate. From this mixture 1 g tablets were compressed to a hardness of 12 kg.

EXAMPLE 17. Toothpaste. A suitable toothpaste base for incorporating the compounds of the invention is described in U.S. Patent No. 3,935,305 (Delaney et al.) which is incorporated herein by reference; in particular, see Example 1. A suitable amount of ArgGlyArgProGln acetate (for preparation, see above) is 1% by weight of base.

EXAMPLE 18. Mouthwash. A suitable mouthwash base for incorporating the compounds of the invention is described in U.S. Patent No. 5,145,664 (Thompson) which is incorporated herein by reference; see Example 1. A suitable amount of ArgGlyArgProGln acetate (for preparation, see above) is 1% by weight of base.

30

LEGENDS TO FIGURES

- Fig 1. Native alkaline PAGE patterns of N-terminal acidic
 peptides from degradation of acidic PRPs (PRP-1 and PRP-3)
 by strains of commensal Streptococcus and Actinomyces
 species. Shown are the degradation patterns of strains

 5 representative for each of the four degradation patterns (IIV) identified among isolates of streptococci and
 actinomycetes (cf. Table 1). All strains (Table 1) were coincubated with purified PRP-1 and PRP-3 for different times
 (15 min, 4h and 20 h) and analyzed for such peptide
 10 patterns.
 - Fig. 2. Gel filtration patterns of fragments formed by degradation of PRP-1 by strains representative for each of the four degradation patterns identified among streptococci and actinomycetes (cf. Table 1). The strains were incubated with PRP-1 for 15 min and 20 h. The retention times of purified PRP-1 and PRP-3 are indicated by vertical arrows.
- Fig 3a. Gel filtration patterns of peptide fragments from degradation of PRP-1 by *S. gordonii* strain SK12 for different times. The numbering of the peaks refer to mass spectrometric identification of the corresponding peptide structures (cf. Fig. 3b). The vertical arrows denote the retention times of purified PRP-1 and PRP-3.

separated using either gel filtration (peaks 1-4, cf. Fig. 3a) or peptide gel filtration (peak 5). The numbering of peptides by letters A-G denotes the corresponding signals in the mass spectrum. Pyr indicates a pyroglutamic acid. The

Fig. 3b. Identification by mass spectrometry of fragments

mass numbers with an asterisk indicate average masses from de-convoluted mass spectra, while unlabeled mass numbers are consistent with the monoisotopic mass of the peptide.

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20 Table 1. Degradation of acidic PRPs by Streptococcus and Actinomyces species

Species ^a	Strain ^a	Acidic I	PRP degr	adation ^b	PRP-1	IgA1
		PRP-1	PRP-3	Type ^c	adhesion ^d	protease ^e
S. gordonii	SK 12, SK 184	5	1	I	-	-
	SK 120	5	0	I	+	-
	SK 33	0	0	-	-	-
S. sanguis	SK 85	4	0	I	-	+
21	SK 112	2	1	II	+	+
	SK 37	2	1	II	-	+
)	SK 162	2	1	III	-	+
S. mitis	SK 304	1	0	II	-	-
	SK 305	0	0	-	+	-
	SK 96, SK 142	0	0	-	-	-
S. anginosus	SK 215	3	1	III	-	-
;	SK 52	1	1	II	-	-
	SK 63	1	0	II	-	-
	SK 218	0	0	-	-	-

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S. oralis	SK 143	1	0	ΙV	-	+
	SK 2, SK 92	0	0	-	-	+
	C 104	0	0	-	nt	+
S. mutans	Ingbritt, JBP	0	0	-	-	nt
	NCTC 10449	0	0	-	nt	nt
S. sobrinus	SL-1	0	0	-	nt	nt
A. odontolyticus	T-5-G	5	5	IV	+	nt
3	T-1-K	2	1	IV	-	nt
	T-23-N,T-3-G	1 ^b	1	IV	-	nt
	T-21-N, T-22-N, PK 984	0	0	-	-	nt
A. naeslundii geno	ospecies 1					
8	ATCC 12104, P-3-N, P-5-N,	0	0	-	-	nt
	P-11-N, B-2-G, PK 947					
A. naeslundii geno	ospecies 2					
Ü	T14V, P-1-N, P-7-N, P-1-K,	0	0	-	+	nt
	P-1-G, B-7-N, P-2-N					
A. viscosus	19246	0	0	-	-	nt
P. gingivalis	W83	5	5	-	-	+
	S. mutans S. sobrinus A. odontolyticus A. naeslundii geno A. naeslundii geno A. viscosus	SK 2, SK 92 C 104 S. mutans Ingbritt, JBP NCTC 10449 S. sobrinus SL-1 A. odontolyticus T-5-G T-1-K T-23-N,T-3-G T-21-N, T-22-N, PK 984 A. naeslundii genospecies 1 ATCC 12104, P-3-N, P-5-N, P-11-N, B-2-G, PK 947 A. naeslundii genospecies 2 T14V, P-1-N, P-7-N, P-1-K, P-1-G, B-7-N, P-2-N A. viscosus 19246	SK 2, SK 92 C 104 S. mutans Ingbritt, JBP NCTC 10449 S. sobrinus SL-1 A. odontolyticus T-5-G T-1-K 2 T-23-N,T-3-G T-21-N, T-22-N, PK 984 A. naeslundii genospecies 1 ATCC 12104, P-3-N, P-5-N, P-11-N, B-2-G, PK 947 A. naeslundii genospecies 2 T14V, P-1-N, P-7-N, P-1-K, P-1-G, B-7-N, P-2-N A. viscosus 19246 0	SK 2, SK 92 0 0 0 C 104 0 0 0 S. mutans Ingbritt, JBP 0 0 0 NCTC 10449 0 0 0 S. sobrinus SL-1 0 0 0 A. odontolyticus T-5-G 5 5 T-1-K 2 1 T-23-N,T-3-G 1 ^b 1 T-21-N, T-22-N, PK 984 0 0 A. naeslundri genospecies 1 ATCC 12104, P-3-N, P-5-N, 0 0 P-11-N, B-2-G, PK 947 A. naeslundri genospecies 2 T14V, P-1-N, P-7-N, P-1-K, 0 0 P-1-G, B-7-N, P-2-N A. viscosus 19246 0 0	SK 2, SK 92 C 104 SK 2, SK 92 C 104 O O O O O S. mutans Ingbritt, JBP NCTC 10449 O O O O O S. sobrinus SL-1 O O O O O O O O O O O O O O O O O O O	SK 2, SK 92 C 104 SK 2, SK 92 C 104 O O - nt S. mutans Ingbritt, JBP O O O - nt S. sobrinus SL-1 A. odontolyticus T-5-G T-1-K T-1-K T-23-N,T-3-G T-21-N, T-22-N, PK 984 A. naeslundii genospecies 1 ATCC 12104, P-3-N, P-5-N, O O - P-11-N, B-2-G, PK 947 A. naeslundii genospecies 2 T14V, P-1-N, P-7-N, P-1-K, O O - + P-1-G, B-7-N, P-2-N A. viscosus 19246 O O O

- 20 a) The sources of the strains are given above.
 - b) Acidic PRP degradation was carried out by co-incubation of PRP-1 or PRP-3 and bacteria for 15 min, 4 h and 20 h followed by native alkaline PAGE (Fig.1). The degree of cleavage at 4 h was scored from 0 to 5 by densitometry: 0 = 0<10 %,</p>
- 1 = 10<20 %, 2 = 20<40 %, 3 = 40<60 %, 4 = 60<80 %, and
 5 = 80<100 % loss of acidic PRPs. Score 1 labeled with a
 superscript denotes the formation of degradation products.

 Cells free supernatant resulted in similar degradation
 patterns for most strains. 95 % confidence intervals of
 densitometric runs corresponded to 8 % of mean.
 - c) The strains were classified into degradation patterns I-IV based on native alkaline PAGE peptide profiles (Fig.1). The type-I pattern displayed a peptide migrating just below PRP-3; The type-II pattern a peptide migrating just below PRP-3 and another between PRP-1 and PRP-3; The type-III pattern two peptides migrating just below and above PRP-3, one between PRP-1 and PRP-3 and a fourth just below PRP-1; and the type-IV pattern a peptide migrating identical to PRP-3 and another between PRP-1 and PRP-3 although closer to PRP-1 compared to patterns II and III.

- d) Adhesion of ³⁵S-labeled bacteria to PRP-1-coated hydroxyapatite beads. Adhesion exceeding 15 % of added cells was considered positive. Adhesion of *Actinomyces* was tested at 5μg/ml of PRP-1 and for other bacteria at 30 μg/ml of PRP-1. The criteria for positive binding and assay conditions differ from those in other studies of adhesion of streptococci and *P. gingivalis* (Hsu et al., 1994).
- e) S-IgA protease activity of commensal streptococci (Kilian et al., 1989), P. gingivalis (Kilian, 1981) and Actinomyces species, which lack S-IgA1 protease activity (M. Kilian, personal communication).

Table 2. Simplified map of generated peptides in comparison with PRP-1.

Peak	Peptide ,		Mass (Da)
1	Pyr1Pro149-Gln150		15 514.0*
2	Pyr1Pro104-Pro105		11 005.5*
3	Gly111Pro149-Gln150		3 930.0
4	Pro96Pro109-Gln110	(A)	1 663.7
4	Gly111Pro130-Gln131	(B)	2 082.1
4	Gly111Pro134-Pro135	(C)	2 430.4
4	Gly111Pro135-Gln136	(D)	2 558.4
4	Gly111Pro140-Gln141		
4	Gly111Pro149-Gln150		
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- 4 Pro104---Pro149-Gln150 (G) 4 718.3 5 Oligopeptides/amino acids
- 5 The potential cleavage sites at peptide bonds formed at Pro or Gln residues are given. The presumed Arg-containing pentapeptide is blown up, ProGln-termini are marked by black circles and the post-translational cyclization of the N-terminal Glu residue to a pyroglutamic acid residue is marked Pyr. The verification by mass spectrometry of phosphorylation of Ser at positions 8 and 22 is marked.

Claims

- An oligopeptide protecting against dental caries
 comprising two arginine residues selected from the group consisting of pentapeptide, hexapeptide, heptapeptide, octapeptide, nonapeptide and decapeptide.
- The oligopeptide of claim 1 comprised by the sequence
 GlyGlyHisProArgProProArgGlyArgProGlnGlyProProGlnGln.
 - 3. The oligopeptide of claim 2 ArgGlyArgProGln.
 - 4. The oligopeptide of claim 2 selected from:
- 15 ArgGlyArgProGln;

ArgGlyArgProGlnGly;

ArgGlyArgProGlnGlyPro;

ArgGlyArgProGlnGlyProPro;

ArgGlyArgProGlnGlyProProGln;

- 20 ArgGlyArgProGlnGlyProProGlnGln;
 - GlyGlyHisProArgProProArgGlyArg;

GlyHisProArgProProArgGlyArg;

HisProArgProProArgGlyArg;

ProArgProProArgGlyArg;

- 25 ArgProProArgGlyArg;
 - ProProArgGlyArg.
 - 5. A method of preventing dental caries comprising the oral administration of a prevention-effective amount of the peptide of any of claims 1-4.
 - 6. A composition for preventing dental caries comprising a prevention-effective amount of the oligopeptide of any of claims 1-4 and a suitable carrier.

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7. The composition of claim 6 wherein the carrier is selected from the group consisting of toothpaste, mouthwash chewing-gum, lozenge, chewable tablet.

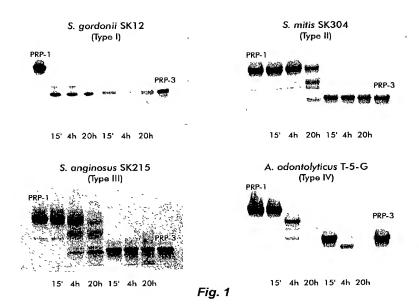
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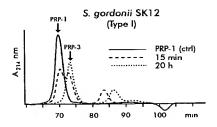
8. The manufacture of a medicament for prevention of dental caries comprising a prevention-effective amount of the oligopeptide of any of claims 1-4.

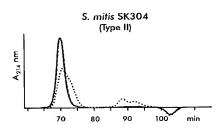
Abstract

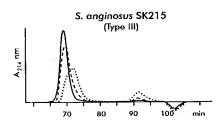
A penta- to decapeptide containing two or more Arg protecting against dental caries is comprised by the amino acid sequence GlyGlyHisProArgProProArgGlyArgProGlnGlyProProGlnGln. Also disclosed is a composition for preventing dental caries comprising an effective amount of the peptide, and a corresponding method of use.

1009749/06/9909









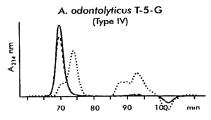


Fig. 2

. Hanning

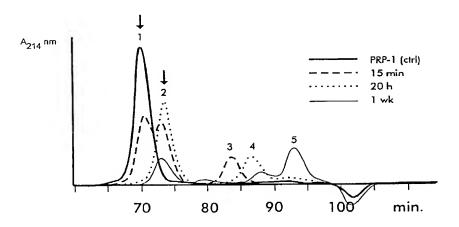


Fig. 3a

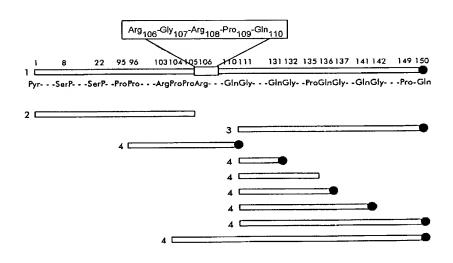


Fig. 3b

							FILE NO. 432.0044		
-	As a below named inventor, I hereby declare that: my residence, post office address and citizenship are as stated below new believe that I am the original, first and sole inventor (if only one name is listed below) or a joint inventor (if plural inventors matter which is claimed and for which a patent is sought on the invention entitled: PREVENTION OF DENTAL CARIES							hat I verily the subject	
	the specification of which is attac was filed on November application number 10/00	patent							
	I hereby state that I have reviewed amendment referred to above. I acknowledge the duty to disclose \$1.56. I hereby claim priority benefits ur States provisional application(s) listed date before that of the application on	e all information known der Title 35, United St below and have also ic	a to be material to parates Code §119 of a dentified below any	tentability in accordar	ice with Ti	itle 37, Co	de of Federal Re	egulations,	
	Prior Foreign or Provisional Applicat		LLUW CDED	5.4572.07			DOLOD WOLL		
	COUNTRY	APPLICATION	NUMBER	DATE OF (day, mon			PRIORITY (UNDER 35 U		
	Sweden	99017	73-3	May 17	, 1999		YES X		
	PCT	PCT/SE00		May 11			YES X		
							YES	NO	
	I hereby claim the benefit under T matter of each of the claims of this ar Title 35, United States Code, §112, I Federal Regulations, §1.56 which bec this application.	itle 35, United States (plication is not disclose acknowledge the duty came available between	code, §120 of any U ed in the prior Unite to disclose informati the filing date of the	nited States application of States application in on which is material to prior application and	n(s) listed the manne o patentab the nation	below and er provided ility as def nal or PCT	, insofar as the s i by the first par ined in Title 37, international fil	ubject agraph of Code of ing date of	
	UNITED STATES APPLICATION NUMBER	UNITED STATES DATE OF FILING						oned)	
	I hereby appoint customer no 321 Reg. No. 24,735, Gary M. Hoffman, Gregory, Reg. No. 28,954, Stephen 2 J. Thronson, Reg. No. 33,082, Micha and revocation to prosecute this applic correspondence.	172 DICKSTEIN, SHA Reg. No. 26,411, Stev A. Soffen, Reg. No. 31 ael J. Scheer, Reg. No. cation, to transact all bu	PIRO, MORIN & Oven I. Weisburd, Re, 063, James W. Bra 34,425, and Eric Ousiness in the Patent	SHINSKY, LLP, and g. No. 27,409, Thoma dy, Jr., Reg. No. 32, diver, Reg. No. 35,30 & Trademark Office	the members J. D'An 115, Jon D 7, as attor connected	pers of the nico, Reg. D. Grossmaneys with therewith	firm, Edward A No. 28,371, Do an, Reg. No. 32, full power of sul and to receive al	. Meilman, nald A. 699, Mark ostitution l	
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٠	RESIDENCE (City and either State or Foreign Country) S-903 37 Umea, Sweden Sweden						ENSHIP 6	SEX	
	post office address Ragangen, 6A, S-903 37	Umea, Swede	n						
-cN	FULL NAME OF SECOND JOINT INVENT INGEGER JOHANSSON	' '	INVENTOR'S SIGNA	TURE		date 2000	r-65 -3	L5	
	RESIDENCE (City and either State or For S-903 39 Umea, Sweden			V	countre Swed	y of citiz en	L ~ 65 ~ 3	SEX	
ļ	Post office address Svampvaagen 18, S-903	39 Umea, Swe	den						

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